

11
5/8/02

1600

R. mitra
RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/667,380A

DATE: 05/02/2002
 TIME: 17:21:26

Input Set : A:\LEX-0042-USA SEQLIST.txt
 Output Set: N:\CRF3\05022002\I667380A.raw

ENTERED

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4 <110> APPLICANT: Donoho, Gregory
5   Turner, C. Alexander Jr.
6   Wattler, Frank
7   Nehls, Michael
8   Friedrich, Glenn
9   Zambrowicz, Brian
10  Sands, Arthur T.
12 <120> TITLE OF INVENTION: Novel Human Protease Inhibitor-Like
13  Proteins and Polynucleotides Encoding the Same
16 <130> FILE REFERENCE: LEX-0042-USA
C--> 18 <140> CURRENT APPLICATION NUMBER: US/09/667,380A
C--> 18 <141> CURRENT FILING DATE: 2000-09-22
18 <150> PRIOR APPLICATION NUMBER: US 60/156,101
19 <151> PRIOR FILING DATE: 1999-09-24
21 <160> NUMBER OF SEQ ID NOS: 3
23 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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26 <211> LENGTH: 1491
27 <212> TYPE: DNA
28 <213> ORGANISM: homo sapiens
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33 aacgagtctc actccgggt ccgcagagcc atccccaggg aggacaaga ggagatcctc 180
34 atgctgcaca acaagcttcg gggccaggtg cagcctcagg cctccaacat ggagtacatg 240
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38 ccctaccga gcgagtgc aa cccctgggtt ccagagaggt gtcggggcc tatgtgcacg 480
39 cactacacac agatagttt ggcaccacc aacaagatcg ttgtgtctgt gaacacctgc 540
40 cggaaagatga ctgtctgggg agaagtttg gagaacgcgg tctactttt ctgcaattat 600
41 tctccaaagg ggaactggat tggagaaggc ccctacaaga atggccggcc ctgctctgag 660
42 tgcaccacca gctatggagg cagctgcagg aacaacttgt gttaccgaga agaaacctac 720
43 actccaaaac ctgaaacggc cgagatgaat gaggtggaaa cggctccat tcctgaagaa 780
44 aaccatgtt ggctccaacc gagggtgtg agacccacca agcccaagaa aacctctgcg 840
45 gtcaactaca tgacccaagt cgtcagatgt gacaccaaga tgaaggacag gtgcaaagg 900
46 tccacgtgt acaaggtaa gtgcccagca ggctgcctga accacaaggc gaagatctt 960
47 ggaagtctgt tctatgaaag ctcgtctagc atatgcccgcc cgcccatcca ctacggatc 1020
48 ctggatgaca agggaggcct ggtggatatac accaggaacg ggaaggccc cttttcggt 1080
49 aagtctgaga gacacggcgt gcagtcctc agcaaataca aaccttcccg ctcattcatg 1140
50 gtgtcaaaag tgaaagtgc ggatttggac tgctacacga ccgttgctca gctgtcccc 1200
51 tttgaaaagc cagcaactca ctgcccaga atccattgtc cggcacactg caaagacgaa 1260
52 ccttcctact gggctccgggt gtttggaaacc aacatctatg cagataccctc aagcatctgc 1320

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53 aagacagctg tgcacgcggg agtcatcagc aacgagagtg ggggtgacgt ggacgtgatg
 54 cccgtggata aaaagaagac ctacgtggc tcgctcagga atggagttca gtctgaaagc
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 59 <212> TYPE: PRT
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 65 Val Cys Gly Ser Gln Gly Tyr Leu Leu Pro Asn Val Thr Leu Leu Glu
 66 20 25 30
 67 Glu Leu Leu Ser Lys Tyr Gln His Asn Glu Ser His Ser Arg Val Arg
 68 35 40 45
 69 Arg Ala Ile Pro Arg Glu Asp Lys Glu Glu Ile Leu Met Leu His Asn
 70 50 55 60
 71 Lys Leu Arg Gly Gln Val Gln Pro Gln Ala Ser Asn Met Glu Tyr Met
 72 65 70 75 80
 73 Thr Trp Asp Asp Glu Leu Glu Lys Ser Ala Ala Ala Trp Ala Ser Gln
 74 85 90 95
 75 Cys Ile Trp Glu His Gly Pro Thr Ser Leu Leu Val Ser Ile Gly Gln
 76 100 105 110
 77 Asn Leu Gly Ala His Trp Gly Arg Tyr Arg Ser Pro Gly Phe His Val
 78 115 120 125
 79 Gln Ser Trp Tyr Asp Glu Val Lys Asp Tyr Thr Tyr Pro Tyr Pro Ser
 80 130 135 140
 81 Glu Cys Asn Pro Trp Cys Pro Glu Arg Cys Ser Gly Pro Met Cys Thr
 82 145 150 155 160
 83 His Tyr Thr Gln Ile Val Trp Ala Thr Thr Asn Lys Ile Gly Cys Ala
 84 165 170 175
 85 Val Asn Thr Cys Arg Lys Met Thr Val Trp Gly Glu Val Trp Glu Asn
 86 180 185 190
 87 Ala Val Tyr Phe Val Cys Asn Tyr Ser Pro Lys Gly Asn Trp Ile Gly
 88 195 200 205
 89 Glu Ala Pro Tyr Lys Asn Gly Arg Pro Cys Ser Glu Cys Pro Pro Ser
 90 210 215 220
 91 Tyr Gly Gly Ser Cys Arg Asn Asn Leu Cys Tyr Arg Glu Glu Thr Tyr
 92 225 230 235 240
 93 Thr Pro Lys Pro Glu Thr Asp Glu Met Asn Glu Val Glu Thr Ala Pro
 94 245 250 255
 95 Ile Pro Glu Glu Asn His Val Trp Leu Gln Pro Arg Val Met Arg Pro
 96 260 265 270
 97 Thr Lys Pro Lys Lys Thr Ser Ala Val Asn Tyr Met Thr Gln Val Val
 98 275 280 285
 99 Arg Cys Asp Thr Lys Met Lys Asp Arg Cys Lys Gly Ser Thr Cys Asn
 100 290 295 300
 101 Arg Tyr Gln Cys Pro Ala Gly Cys Leu Asn His Lys Ala Lys Ile Phe
 102 305 310 315 320
 103 Gly Ser Leu Phe Tyr Glu Ser Ser Ser Ile Cys Arg Ala Ala Ile

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	325	330	335
104			
105	His Tyr Gly Ile Leu Asp Asp Lys	Gly Gly Leu Val Asp	Ile Thr Arg
106	340	345	350
107	Asn Gly Lys Val Pro Phe Phe Val	Lys Ser Glu Arg	His Gly Val Gln
108	355	360	365
109	Ser Leu Ser Lys Tyr Lys Pro Ser Ser Ser	Phe Met Val Ser Lys Val	
110	370	375	380
111	Lys Val Gln Asp Leu Asp Cys Tyr Thr Val	Ala Gln Leu Cys Pro	
112	385	390	395
113	Phe Glu Lys Pro Ala Thr His Cys Pro Arg	Ile His Cys Pro Ala His	
114	405	410	415
115	Cys Lys Asp Glu Pro Ser Tyr Trp	Ala Pro Val Phe Gly Thr Asn Ile	
116	420	425	430
117	Tyr Ala Asp Thr Ser Ser Ile Cys Lys Thr Ala Val	His Ala Gly Val	
118	435	440	445
119	Ile Ser Asn Glu Ser Gly Gly Asp Val Asp Val	Met Pro Val Asp Lys	
120	450	455	460
121	Lys Lys Thr Tyr Val Gly Ser Leu Arg Asn Gly Val	Gln Ser Glu Ser	
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125 Gln

128 <210> SEQ ID NO: 3

129 <211> LENGTH: 2272

130 <212> TYPE: DNA

131 <213> ORGANISM: homo sapiens

133 <400> SEQUENCE: 3			
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135 tgtgcccgcg ctgtcgccgc tgctaccgcg tctgtggac	gcggggagacg	ccagcgagct	120
136 ggtgatttggga gcccgtcgga gagctcaagc gcccagctct	gccccaggag	cccaaggctgc	180
137 cccgtgagtc ccatagttgc tgcaggatgt gaggcatgag	ctgcgtctgc	ggtgtgtca	240
138 tcccctttggg gctgtgttc ctggctctgc gatcccaagg	ctacctctgc	ccaaacgtca	300
139 ctctctttaga ggagctgctc agcaaataacc agcacacaacga	gtctcactcc	cgggtccgca	360
140 gagccatccc cagggaggac aaggaggaga tcctcatgt	gcacaacaag	tttgggggcc	420
141 aggtgcagcc tcaggcctcc aacatggagt acatgacctg	ggatgacaa	ctggagaagt	480
142 ctgctgcagc gtggggccagt cagtgcacatc gggagcacgg	gcccaccagt	ctgtgggtgt	540
143 ccatcgggca gaacctgggc gctcacttgg gcaggttatcg	ctctccgggg	tccatgtgc	600
144 agtcctgtta tgacgagggt aaggactaca cctaccctta	cccgagcgag	tcaacccct	660
145 ggtgtccaga gaggtgctcg gggcctatgt gcacgcacta	cacacagatc	gtttgggcca	720
146 ccaccaacaa gatcggttgt gctgtgaaca cctgccggaa	gatgactgtc	tggggagaag	780
147 tttggggagaa cgcggtctac tttgtctca attattctcc	aaaggggaaac	tggattggag	840
148 aagccccccta caagaatggc cggccctgtct ctgagtgccc	acccagctat	ggagggcagct	900
149 gcaggaacaa ctttgtttac cgagaagaaa cctacactcc	aaaacctgaa	acggacgaga	960
150 tgaatgaggt gaaaaacggct cccattctg aagaaaaacca	tgtttggctc	caaccgaggg	1020
151 tcatgagacc caccacggccc aagaaaaaccc ctgcggtaaa	ctacatgacc	caagtcgtca	1080
152 gatgtgacac caagatgaag gacaggtgca aagggtccac	gtgtacagg	taccagtgtcc	1140
153 cagcaggctg cctgaaccac aaggcgaaga tctttggaa	tctgttctat	gaaagctcg	1200
154 ctagcatatg cccgcggcc atccactacg ggatcctggaa	tgacaaggaa	ggcctgggtgg	1260
155 atatcaccag gaacgggaag gtcccctct tcgtgaagtc	tgagagacac	ggcgtgcagt	1320

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156 ccctcagcaa atacaaacct tccagctcat tcatgggtgc aaaagtgaaa gtgcaggatt	1380
157 tggactgcta cacgaccgtt gctcagctgt gcccggttga aaagccagca actcaactgcc	1440
158 caagaatcca ttgtccggca cactgcaaag acgaacacctc ctactgggtc ccgggtttg	1500
159 gaaccaacat ctatgcagat acctcaagaca tctgcaagac agctgtgcac gcccggagtca	1560
160 tcagcaacga gagtgggggt gacgtggacg tgatgcccgt ggataaaaaag aagacctacg	1620
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162 cttccggat ctttgcgtc aggcaagtggaa ttccagcac caggggagaa ggggcgtctt	1740
163 caggagggct tcgggggttt gcttttattt ttattttgtc attgcggggt atatggagag	1800
164 tcaggaaact tccttgact gatgttcagt gtccatcaact ttgtggcctg tgggtgaggt	1860
165 gacatctcat cccctcaactg aagcaacagc atcccaaggt gtcagccgg actccctgg	1920
166 gcctgtatcct gctggggcct gggggctcc atctggacgt cctctctctt ttagagatct	1980
167 gagctgtctc ttaaaggggaa cagttgcca aaatgttcct tgctatgtgt tcttctgttgc	2040
168 gtggaggaag ttgatttcaa cccccctgcc aaaagaacaa accatttggaa gtcacaatt	2100
169 gtgaagcatt cacggcgtcg gaagaggcct tttgagcaag cgccaatgag tttaggaaat	2160
170 gaagttagaaat gtagtttattt aaaaataaaa aacacagtcc gtccctacca atagaggaaa	2220
171 atggtttaa tgtttgcgttccagacagac aaatgggtca gagtaagaag gc	2272

VERIFICATION SUMMARY

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Input Set : A:\LEX-0042-USA SEQLIST.txt

Output Set: N:\CRF3\05022002\I667380A.raw

L:18 M:270 C: Current Application Number differs, Replaced Current Application No

L:18 M:271 C: Current Filing Date differs, Replaced Current Filing Date